

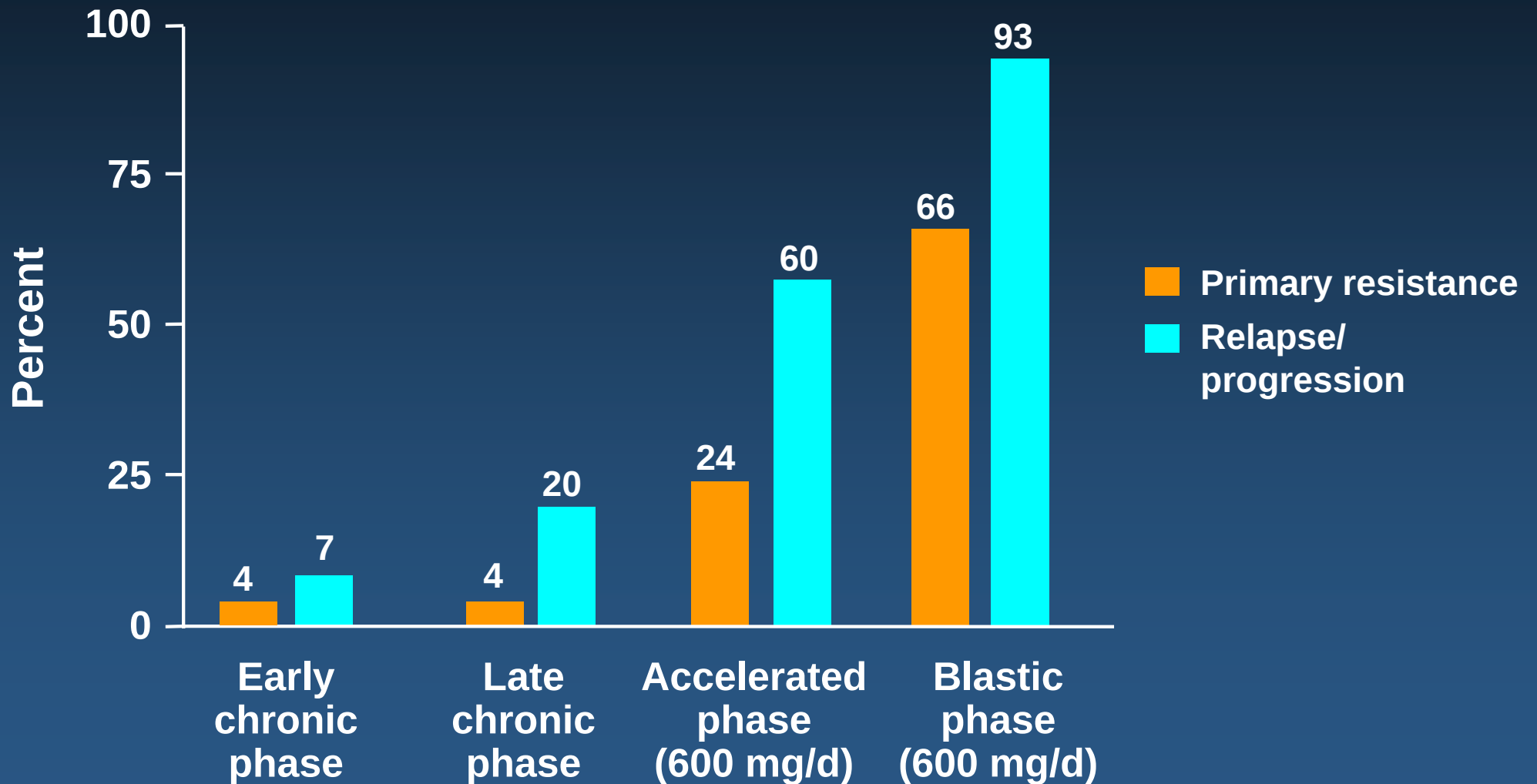


Bcr-Abl Kinase Domain Mutations

Martin Müller



Frequency of imatinib resistance within 3 years



Observations associated with resistance

Genomic amplification / overexpression

BCR-ABL Mutations

Clonal evolution / aneuploidy

Pharmacologic mechanisms

- Association with AGP-levels?

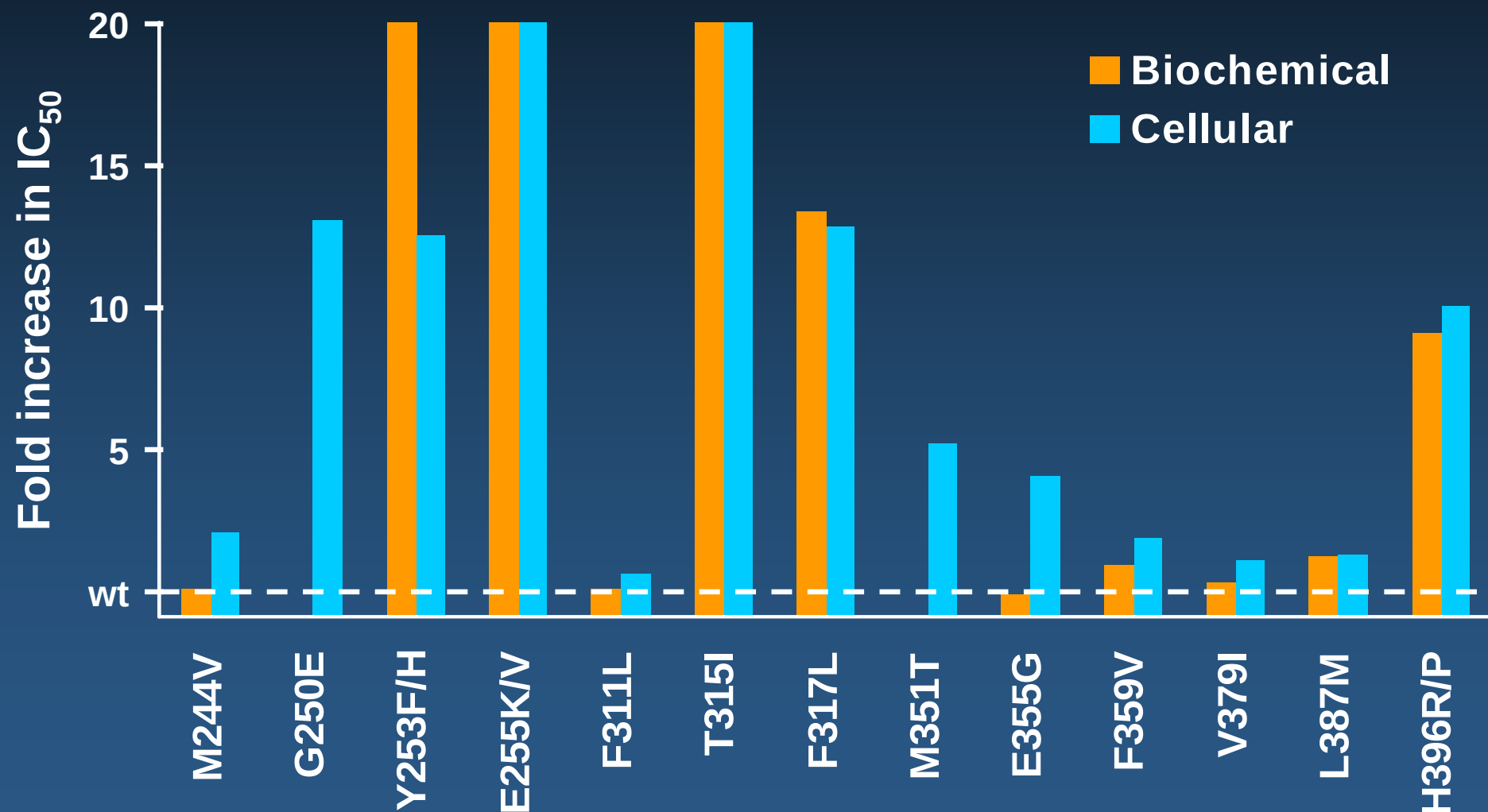
- Low MRP-1 levels independent prognostic factor (Lange et al.)

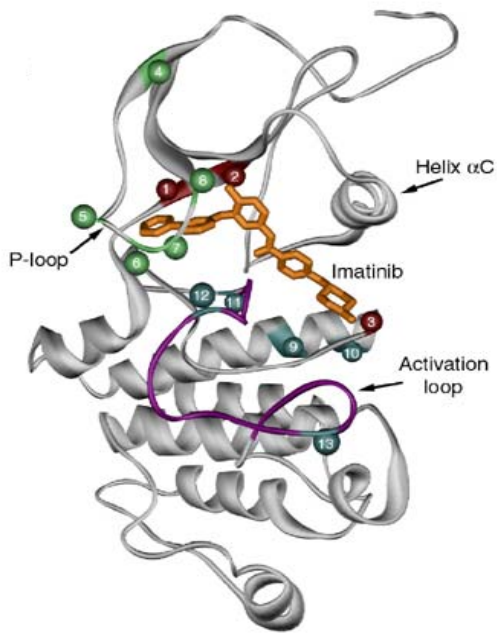
- Decrease of intracellular imatinib-levels in PGP+ patients (Illmer et al.)

Molecular - cytogenetic causes of resistance

Patients With Hematologic Resistance/Relapse	Chronic phase (n=35)	Accelerated phase (n=33)	Blastic phase (n=66)	All (n=134)
<i>BCR-ABL</i> mutations (%)	10/20 (50)	13/21 (62)	10/33 (30)	33/74 (45)
Clonal evolution (%)	15/29 (52)	8/16 (50)	16/22 (73)	39/67 (58)
Combination (%)	5/17 (29)	2/9 (22)	4/17 (24)	11/43 (26)

Mutant BCR-ABL have increased IC_{50} values for imatinib mesylate





Resistance to imatinib: 31 BCR-ABL mutations in 25 amino acids (n=126)

P-Loop

Kinase domain

A-Loop

M244V	L248V	D267G	T315I	M351T	L387F/M	E450G/K
	G250E	T277A	F317L	E355G	M388L	E459K
	Q252H	E279K	L324Q	F359C/V	H396R	
	Y253F/H	F311L	A344V	V379I	A397P	
	E255K/V		A350V			

Polymorphisms: K247R
T315T
E499E

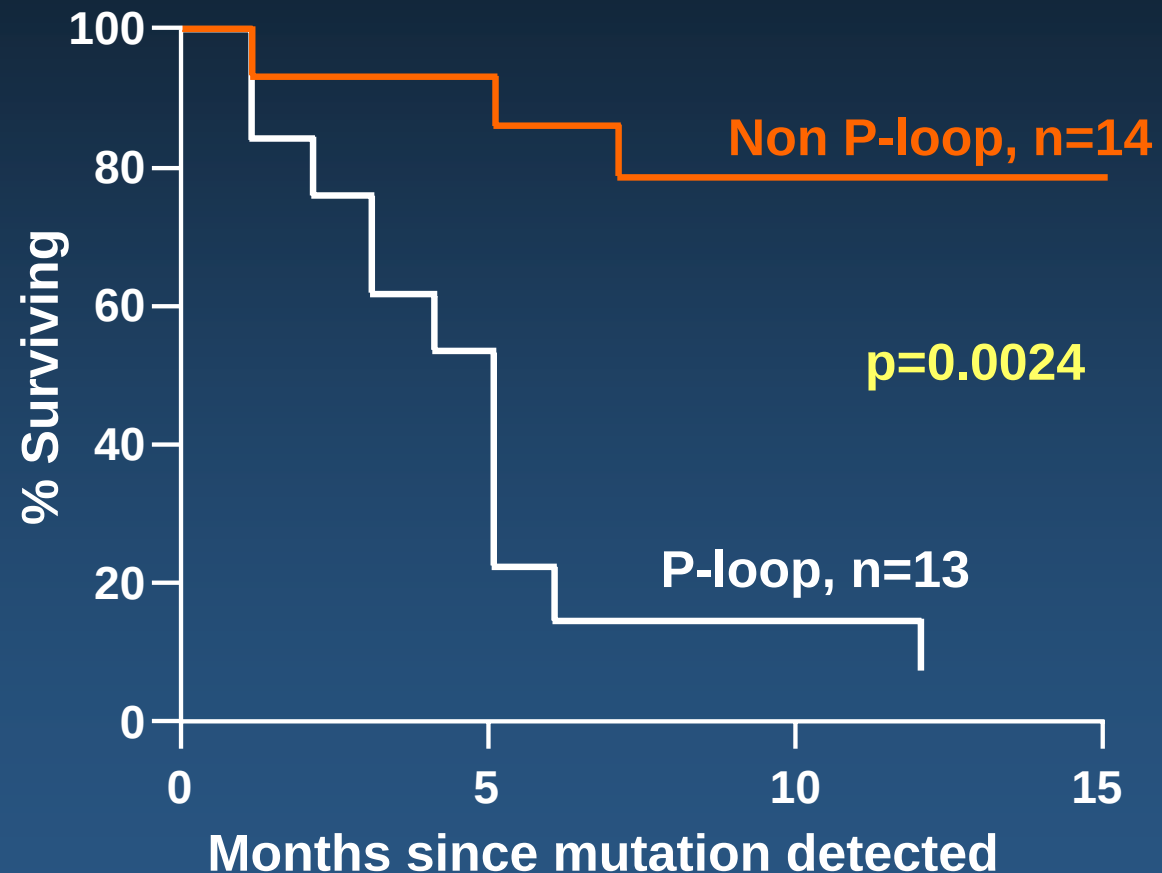
Survival after imatinib failure

Initial Australian experience

n=27

P-Loop: Highly conserved area

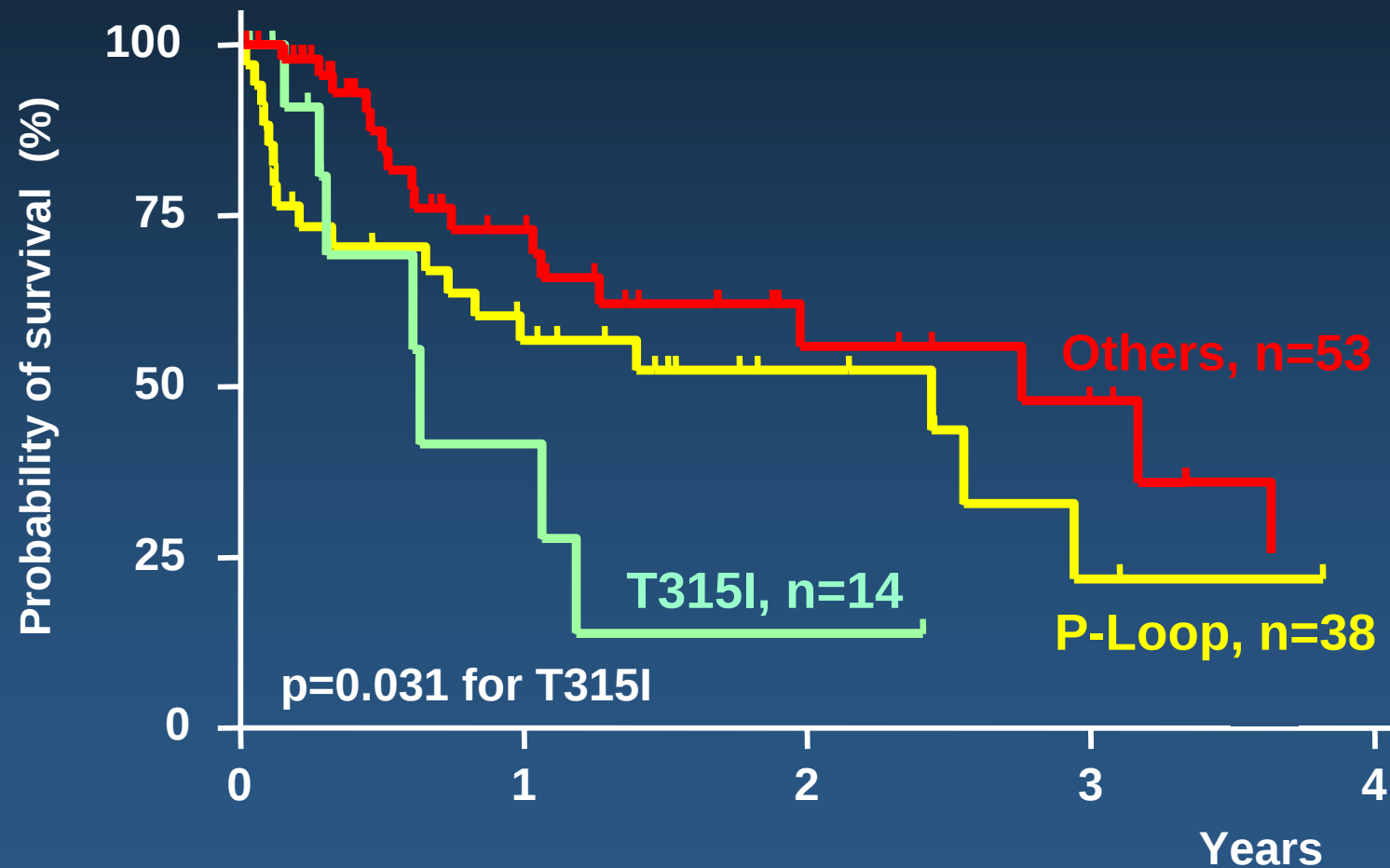
250	252	253	255
G	Q	Y	E
G	H	F	K
G		H	V



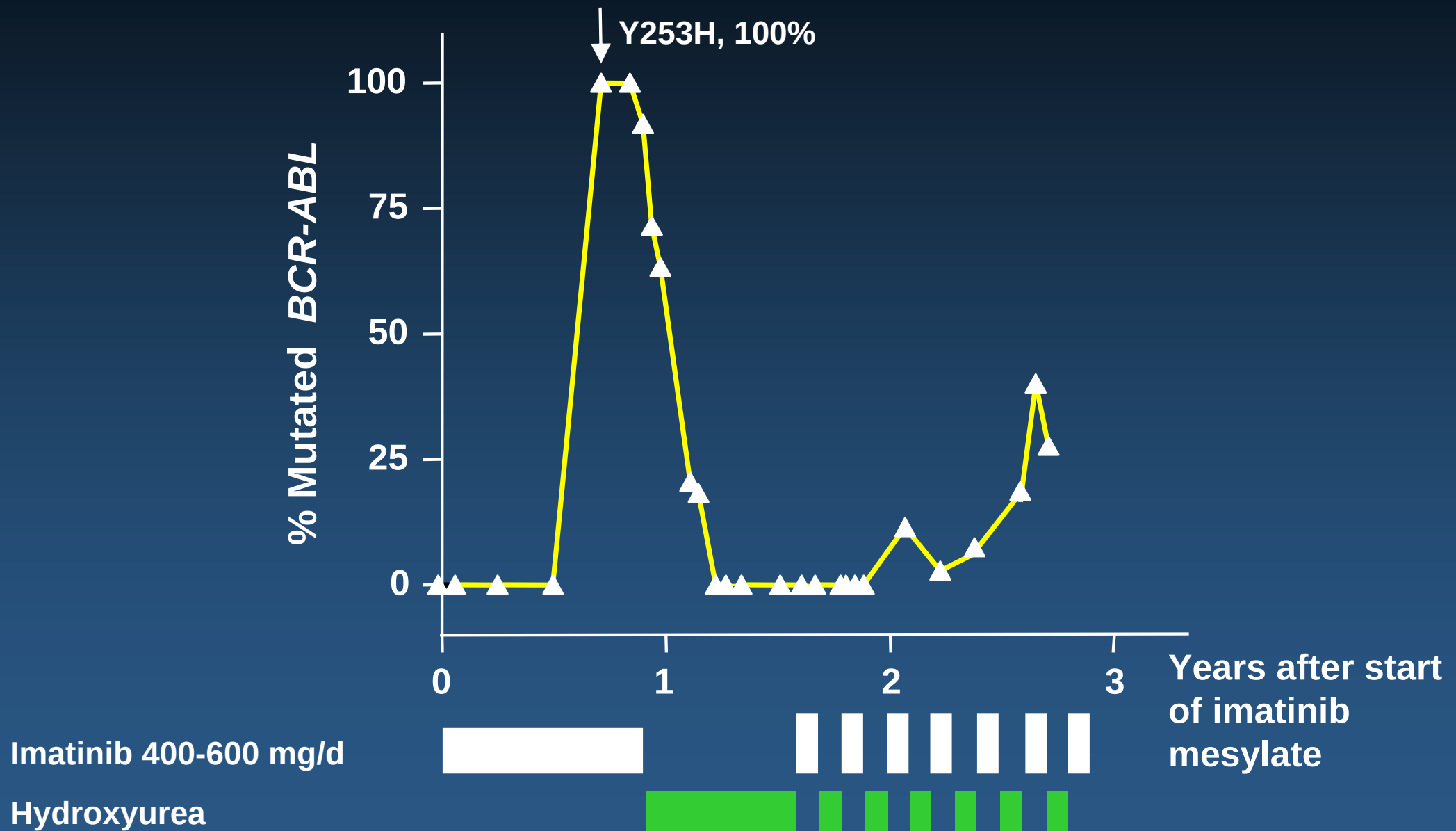
Survival after imatinib failure

Current experience

n=105



CML, chronic phase, IFN-resistant



Significance of P-loop mutations in CML

- 89 imatinib resistant pts in 5 French centers;
26 with P-loop mutations,
18 with T315I, 50 with other mutations
→ significantly worse survival *from diagnosis*
for P-loop or T315I mutations ($p=0.014$)
- 159 pts from MDACC:
52 mutations in 49 pts; 19 P-loop mutations
median F/U 6 mo:
1/19 P-loop vs 4/30 non P-loop died;
15 mo surv 95% vs 83%

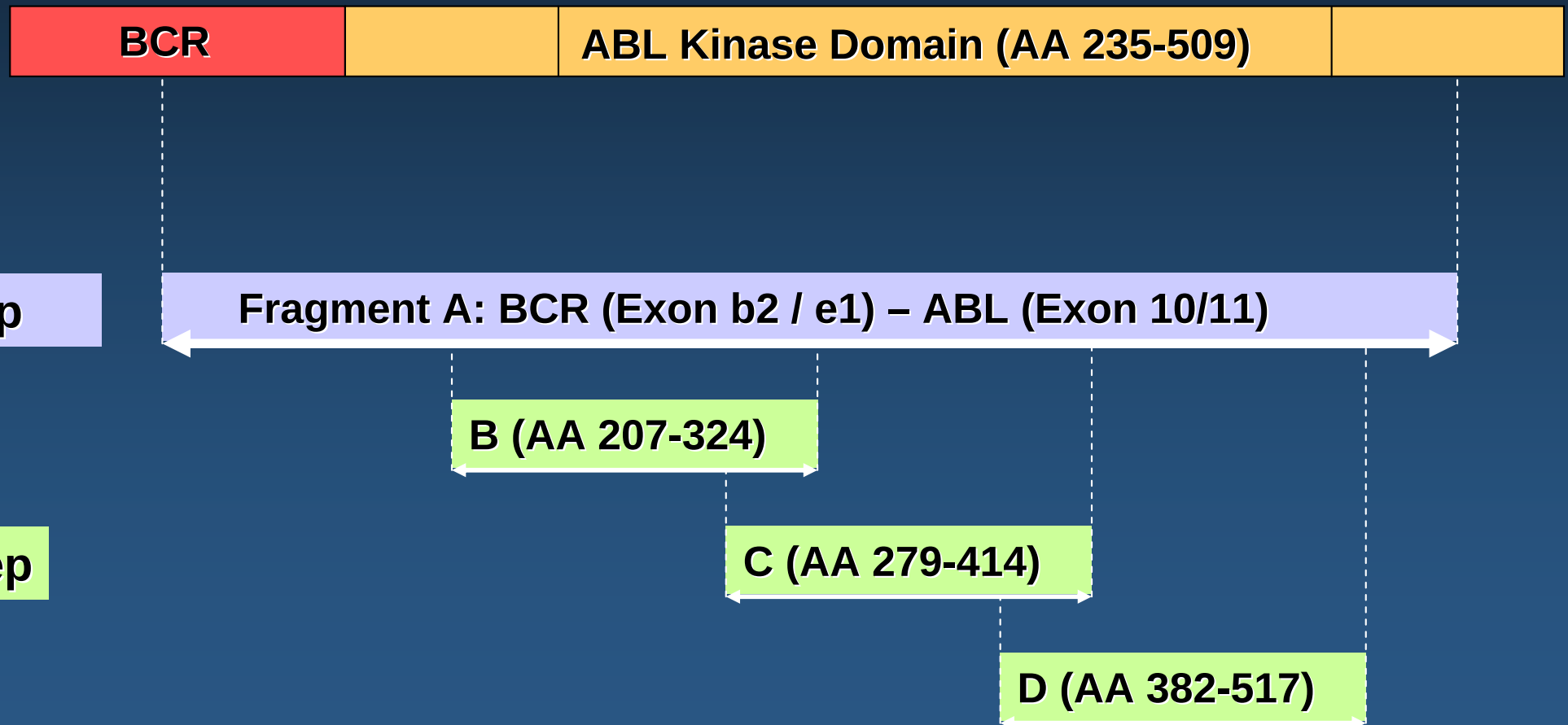
Indications for mutation screening

- hematologic resistance / relapse
- cytogenetic resistance / relapse
- 5-10-fold increase of BCR-ABL load
- prior to therapy with alternative kinase inhibitors (nilotinib, dasatinib)
- 3-monthly intervals under therapy with nilotinib and dasatinib

Methods to detect and quantify BCR-ABL mutations

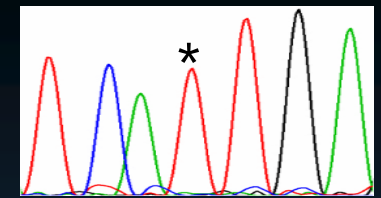
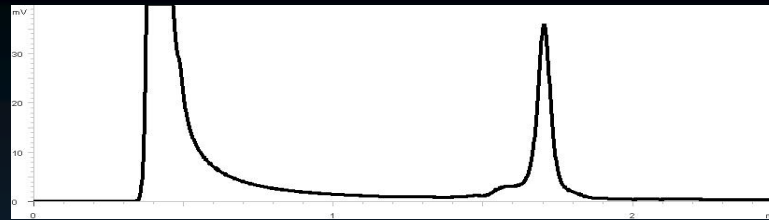
	Specificity	Sensitivity
Sequencing	unspecific	10-20%
Restriction digest analysis	specific	~2-5%
D-HPLC	unspecific	0.1-10%
Allele specific PCR	specific	0.01%
Sequencing of clones	unspecific	1-5%

D-HPLC: nested RT-PCR

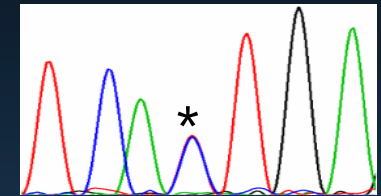
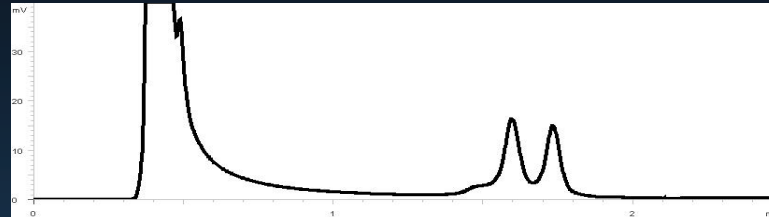


mutant BaF3^{T315I}

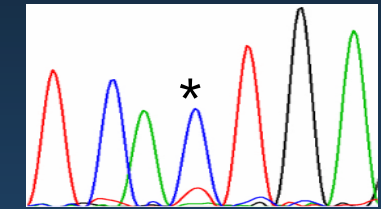
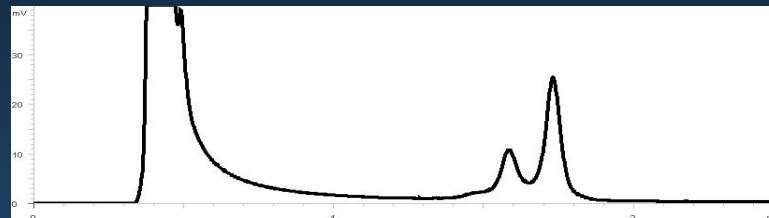
100%



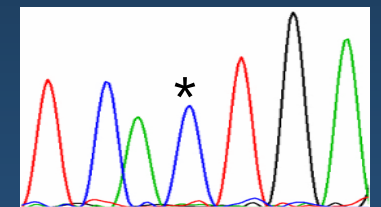
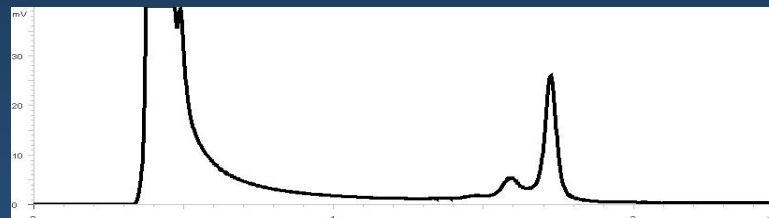
50%



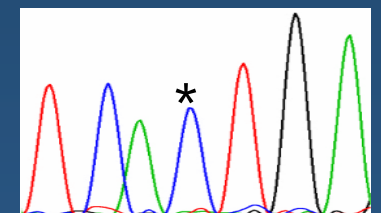
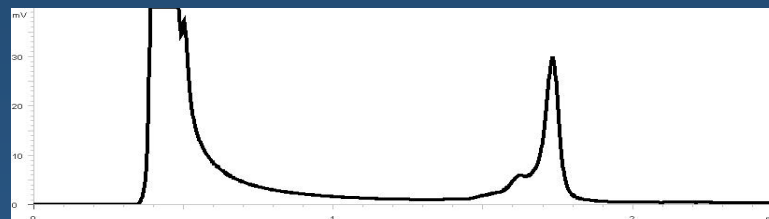
10%



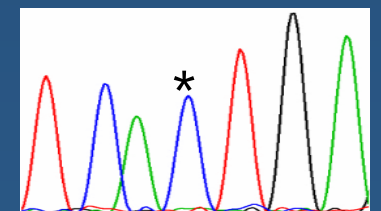
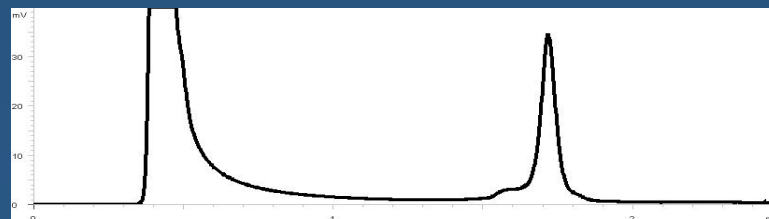
1%



0.1%



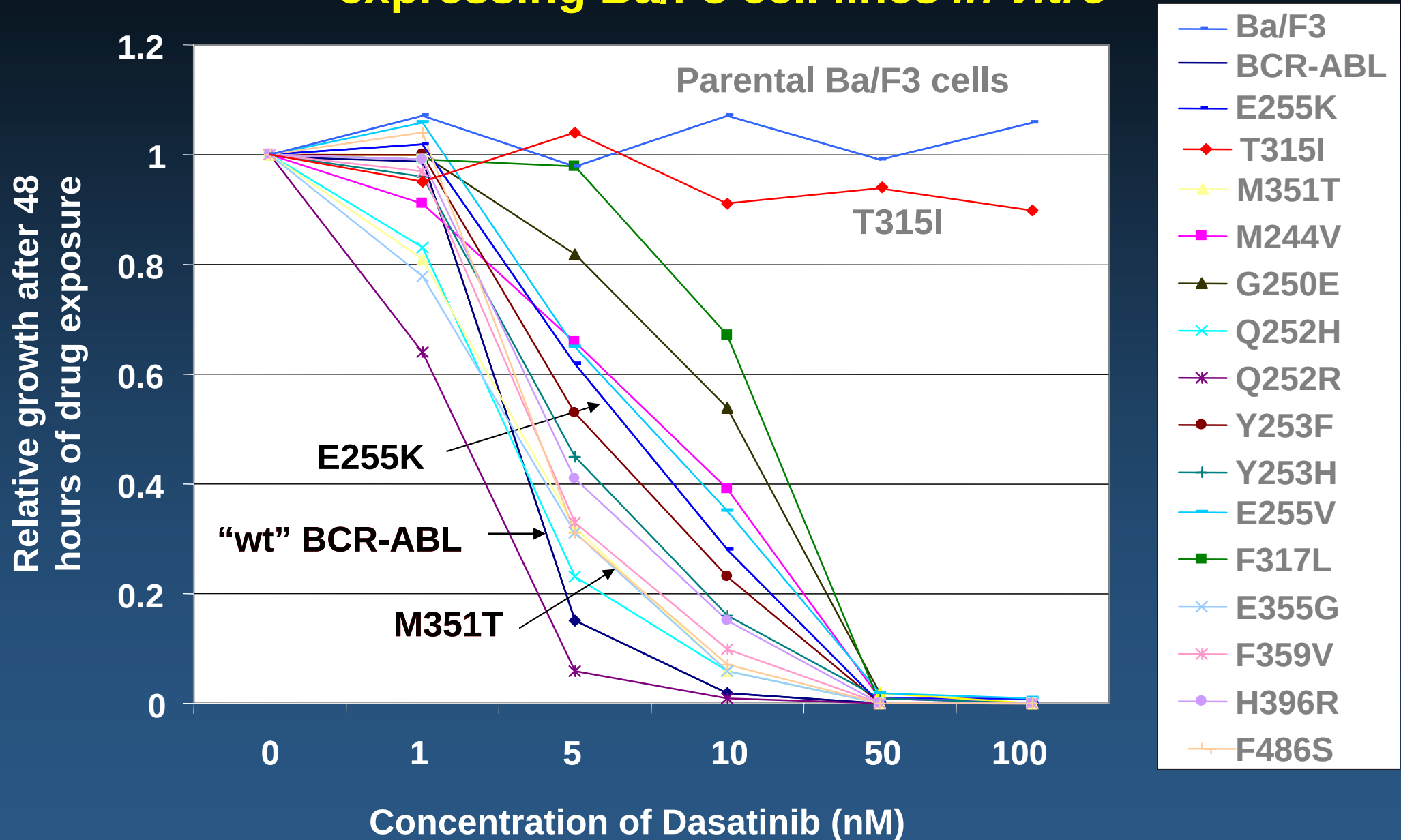
normal BaF3^{BCR-ABL}



Sensitivity of
D-HPLC

0.1-1%

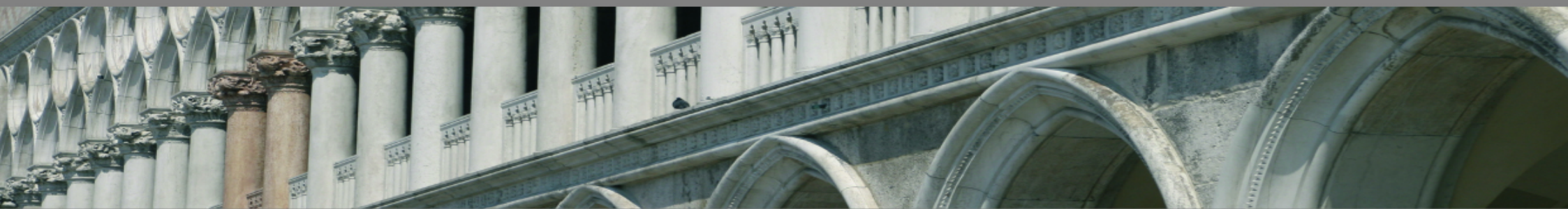
Dasatinib inhibits growth of 14/15 imatinib-resistant BCR-ABL-expressing Ba/F3 cell lines *in vitro*



Conclusions

- Frequency of imatinib resistance depends on the stage of CML.
- The major causes of resistance are BCR-ABL mutations and clonal evolution.
- The impact of early detection of mutated clones by sensitive assays requires prospective evaluation.
- Certain BCR-ABL mutations might especially impair prognosis, in particular on continuous imatinib therapy.
- Early elucidation of imminent resistance to kinase inhibitors might contribute to individualized therapy according to molecular data.

EVOLVING CONCEPTS IN THE MANAGEMENT OF CHRONIC MYELOID LEUKEMIA



RECOMMENDATIONS FROM AN EXPERT PANEL ON BEHALF OF THE EUROPEAN LEUKEMIANET